

(FILE 'HOME' ENTERED AT 12:55:59 ON 12 DEC 2003)

FILE 'MEDLINE, BIOSIS, CAPLUS' ENTERED AT 12:56:09 ON 12 DEC 2003

L1 11 SEA PLU=ON D2S307
L2 4 DUP REM L1 (7 DUPLICATES REMOVED)
D TI 1-4
D IBIB AB 1-4

FILE 'STNGUIDE' ENTERED AT 12:57:50 ON 12 DEC 2003

FILE 'MEDLINE, BIOSIS, CAPLUS' ENTERED AT 13:01:11 ON 12 DEC 2003

L3 59 SEA PLU=ON D2S3346 OR GGAA19E07 OR D2S72 OR D2S105 OR
GATA52A04 OR D2S311 OR D2S1384 OR D2S115
L4 22 DUP REM L3 (37 DUPLICATES REMOVED)
L5 18 SEA PLU=ON L4 NOT L1
D TI 1-18
D IBIB AB 17 18 3 1

FILE 'GENBANK' ENTERED AT 13:05:12 ON 12 DEC 2003

SET NOTICE DISPLAY 1
D ACC X54578 ALL
SET NOTICE LOGIN DISPLAY

FILE 'MEDLINE, BIOSIS, CAPLUS' ENTERED AT 13:05:55 ON 12 DEC 2003

L6 6 SEA PLU=ON L5 AND PY<1999
D TI 1-5
D IBIB AB 1-5

FILE 'STNGUIDE' ENTERED AT 13:08:27 ON 12 DEC 2003

FILE 'MEDLINE, BIOSIS, CAPLUS' ENTERED AT 13:11:17 ON 12 DEC 2003

L7 0 SEA PLU=ON 2Q33 AND (DS2?)
L8 76 SEA PLU=ON 2Q33 AND (D2S?)
L9 20 SEA PLU=ON L8 AND PY<1999
L10 7 DUP REM L9 (13 DUPLICATES REMOVED)
D TI 1-7

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OM nucleic - nucleic search, using sw model

Run on: October 22, 2003, 22:00:05 ; Search time 454.133 Seconds
(without alignments)
1801.657 Million cell updates/sec

Title: US-10-085-906-352

Perfect score: 20

Sequence: 1 tgcactccagcctgagcgac 20

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2888711 seqs, 20454813386 residues

Total number of hits satisfying chosen parameters: 5777422

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*

identical search repeated in GENSEQ, USPATS, USPGPUBS, and EST databases

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Run on: October 22, 2003, 22:00:05 ; Search time 454.133 Seconds
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Title: US-10-085-906-353

Perfect score: 20

Sequence: 1 ttcaacacttaagaatgggg 20

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2888711 seqs, 20454813386 residues

Total number of hits satisfying chosen parameters: 5777422

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identical search repeated in GENSEQ, USPATS, USPGPUBS, and EST databases

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OM nucleic - nucleic search, using sw model

Run on: October 22, 2003, 22:00:05 ; Search time 794.733 Seconds
(without alignments)
1801.657 Million cell updates/sec

Title: US-10-085-906-354

Perfect score: 35

Sequence: 1 attttatttttatttttatttttatttttattttt 35

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2888711 seqs, 20454813386 residues

Total number of hits satisfying chosen parameters: 5777422

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*

identical search repeated in GENSEQ, USPATS, USPGPUBS, and EST databases